



RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number: 101517,039
Source: PCT
Date Processed by STIC: 12-22-04

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
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Revised 05/17/04

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/517,039

DATE: 12/22/2004
TIME: 09:15:16

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Output Set: N:\CRF4\12222004\J517039.raw

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6 <120> TITLE OF INVENTION: METHOD OF TRANSFORMING SOYBEAN
8 <130> FILE REFERENCE: 70094 USPS
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/517,039
C--> 10 <141> CURRENT FILING DATE: 2004-12-07
10 <160> NUMBER OF SEQ ID NOS: 4
12 <170> SOFTWARE: PatentIn version 3.2

ERRORED SEQUENCES

Does Not Comply
Corrected Diskette Needed
(pg. 1-15)

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RAW SEQUENCE LISTING
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386 <223> OTHER INFORMATION: pNov2147

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E--> 458 caaatattcT gaaatgAGCT gttgacaATT aatcatccgg ctcgtataAT gtgtgGAATT 1920
E--> 460 gtgagcggat aacaatttca cacaggAAAC agaccatgag ggaagcgtt atgcggAAAG 1980
E--> 462 tatacgactca actatcAGAG gtatTTggcG tcatacgatc ccatctcgAA ccgacgTTc 2040
E--> 464 tggccgtaca tttgtacggc tccgcagtgg atggcggcCT gaagccACAC agtGatATTG 2100
E--> 466 atttgcTggT tacggTgacc gtaaggCTT atgaaACAC gcccgcAGCT ttGatcaACG 2160
E--> 468 acctttggA aacttcggCT tcccctggAG agagcgaGAT tctccgcgt gtAgAAAGTC 2220
E--> 470 ccattgttGt gcacgacgac atcattccgt ggcgttATCC agctaAGCGC gaactgcaAT 2280
E--> 472 ttggagaATG gcagcgaAT gacattctg caggatCTT cgcAGCCAC acgatcgaca 2340
E--> 474 ttgatctggc tatctgctg acaAAAGCAA gagaACATAG cgttgcTTG gtaggtccAG 2400
E--> 476 cggcggagGA actctttgat cggTTcCTG aacaggatCTT atttgaggcG ctaaatgAAA 2460

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/517,039

DATE: 12/22/2004
TIME: 09:15:16

Input Set : A:\Sequence Listing-70094USPCT.txt
Output Set: N:\CRF4\12222004\J517039.raw

E--> 478 ccttaacgct atggaaactcg ccggcccgact gggctggcga tgagcgaat gtagtgccta 2520
E--> 480 cgttgtcccc cattttggtag acgcgcgatggaa ccggcaaaat cgcgcggaaag gatgtcgctg 2580
E--> 482 ccgactgggc aatggagcgc ctgcggcccc agtatacgcc cgtcataactt gaagctagggc 2640
E--> 484 aggcttatct tggacaagaagaa gatcgcttgg cgtcgccgcgca agatcagttg gaagaatttg 2700

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W--> 489 70094 Syngenta

E--> 491 ttcactacgt gaaaggcgag atcacccaaag tagtcggcaa ataaagctct agtggatctc 2760
E--> 493 cgtacccccc ggggatctgg ctcgcggcgg acgcacgacg ccggggcgag accataggcg 2820
E--> 495 atctcctaaa tcaatagtag ctgtAACCTC gaagcgttcc acttgtaaca acgattgaga 2880
E--> 497 attttgtca taaaattgaa atacccgtt cgcatttttgc tcatccgcgg tcagcccaa 2940
E--> 499 ttctgacgaa ctgcccattt agctggagat gattgtacat cttcacgtg aaaatttctc 3000
E--> 501 aagcgctgtg aacaagggtt cagatttttag attgaaaggt gagccgttga aacacgttct 3060
E--> 503 tcttgcgtat gacgacgtcg ctatgcggca tcttatttattt gaatacctt aCGATCCACGC 3120
E--> 505 cttcaaaagtg accgcggtag ccgacagcac ccagttcaca agagtaactct cttccgcgac 3180
E--> 507 ggtcgatgtc gtgggttgcg atctaaattt aggtcgtaa gatgggctcg agatcggtcg 3240
E--> 509 taatctggcg gcaaagtctg atattccaaat cataattatc agtggcgacc gccttgagga 3300
E--> 511 gacggataaaa gttgttgcac tcgagctagg agcaagtgtat ttatcgcta agccgtttag 3360
E--> 513 tatcagagag tttctagcac gcattcgggt tgccttgcgc gtgcgcccca acgttgcgc 3420
E--> 515 ctccaaagac cgacggcttt tttgttttac tgactggaca cttaatctca ggcaacgtcg 3480
E--> 517 ctgtatgtcc gaagctggcg gtgaggtgaa acttacggca ggtgagttca atcttctcct 3540
E--> 519 cgcgtttta gagaaacccc gcgacgttct atcgcgcgag caacttctca ttgccagtcg 3600
E--> 521 agtacgcgac gaggagggtt atgacaggag tatagatgtt ctcatggta ggctgcgc 3660
E--> 523 caaacttgag gcagatccgt caagccctca actgataaaa acagcaagag gtgcgggta 3720
E--> 525 tttcttgac gcggacgtgc aggttcgca cggggggacg atggcagcct gagccaattc 3780
E--> 527 ccagatcccc gaggaatcgg cgtgagcggc cgcaaacat cggccccggg acaaatacg 3840
E--> 529 gcggcgctgg gtgatgaccc ggtggagaag ttgaaggccg cgcaggccgc ccagcggcaa 3900
E--> 531 cgcatcgagg cagaagcacg ccccggtgaa tcgtggcaag cggccgctga tcgaatccgc 3960
E--> 533 aaagaatccc ggcaaccggc ggcagccggc ggcgcgtcga ttaggaagcc gccaagggc 4020
E--> 535 gacgagcaac cagatttttt cgtttcgatg ctctatgacg tgggcaccccg cgatagtcg 4080
E--> 537 agcatccatgg acgtggccgt tttccgtctg tcgaagcgtg accgacgacg tggcgagggt 4140
E--> 539 atccgctacg agcttccaga cgggcacgtc gaggtttccg caggggccggc cggcatggcc 4200
E--> 541 agtgtgtggg attacgaccc ggtactgtatgcgatgggttttttcc atctaaccga atccatgaaac 4260
E--> 543 cgataccggg aagggaaggg agacaagccc ggccgcgtgt tccgtccaca cgttgcggac 4320
E--> 545 gtactcaagt tctgcccggc agccatggc gggaaagcaga aagacgaccc ggtagaaacc 4380
E--> 547 tgcattcgggt taaacaccac gcacgttgcg atgcagcgtt cgaagaaggc caagaacggc 4440
E--> 549 cgcctgggtga cggtatccga ggggtgaagcc ttgattagcc gctacaagat cgtaaagaggc 4500
E--> 551 gaaaccgggc ggccggagta catcgagatc gagctagctg attggatgtt cccggagatc 4560
E--> 553 acagaqqca aqaaacccqqa cgtqctgacq qttcacccq attactttt qatcgatccc 4620

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W--> 557 70094 Syngenta

E--> 559 ggcatcgccc gttttctcta ccgcctggca cgccgcggc caggcaaggc agaagccaga 4680
E--> 561 tggttgttca agacgatcta cgaacgcagt ggcagcggc gagagttcaa gaagttctgt 4740
E--> 563 ttcaccgtgc gcaagctgat cgggtcaaat gacctgcgg agtacgatt gaaggaggag 4800
E--> 565 gccccggcagg ctggcccgat cctagtcatt cgctaccgca acctgatcga gggcgaagca 4860
E--> 567 tccgcgcgtt cctaattgtac ggagccatg cttagggcaaa ttgccttagc aggggaaaaaa 4920
E--> 569 ggtcgaaaaag gtctctttcc tgtggatagc acgtacattt ggaacccaaa gccgtacatt 4980
E--> 571 gggaaaccgga acccgatcat tggaaaccca aagccgtaca ttggaaaccg gtcacacatg 5040
E--> 573 taagtgtactg atataaaaaga gaaaaaaggc gatttttccg cctaaaaactc tttaaaactt 5100
E--> 575 attaaaaactc taaaaaccccg cctggcctgt gcataactgt ctggccagcg cacagccgaa 5160

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/517,039

DATE: 12/22/2004
TIME: 09:15:16

Input Set : A:\Sequence Listing-70094USPCT.txt
Output Set: N:\CRF4\12222004\J517039.raw

E--> 577 gagctgcaaa aagccctac cttcggtcg ctgcgtccc tacgccccgc cggttgcgt 5220
 E--> 579 cggcctatcg cggccgtgg ccgctcaaaa atggctggcc tacggccagg caatctacca 5280
 E--> 581 gggcgccgac aagccgcgcc gtcgccactc gacggccggc gctgaggctc gcctcgtaa 5340
 E--> 583 gaagggtttg ctgactcata ccaggcctga atgcceccat catccagcca gaaagtggagg 5400
 E--> 585 gagccacgg tgatgagagc tttgttga⁹ gtggaccagt tggatgtttt gaacttttgc 5460-5459
 E--> 587 ttgccacgg aacggcttcg gttgtcgga agatcggtga tctgtatcctt caactcagca 5520
 E--> 589 aaagttcgat ttattcaaca aagccgcgt cccgtcaagt caggtataatg ctctgcagg 5580-5579
 E--> 591 gttacaacca attaaccat tctgatttga aaaactcata gggatcaaaa taaaacttgca 5640
 E--> 593 attattcat atcaggatta tcaataccat atttt⁹aaa aagccgttcc tggatgtttt 5700-5699
 E--> 595 gagaaaactc accgaggcag ttccatagga tggcaagatc ctgtatcgg tctgcgattc 5760
 E--> 597 cgactcgtcc aacatcaata caacctatta atttccctc gtcaaaaata aggttatcaa 5820
 E--> 599 gtgagaaatc accatgagtg acgactgaat ccggtagaa tggcaaaaage tctgcattaa 5880
 E--> 601 tgaatcgcc aacgcgcggg gagaggccgt ttgcgtattt ggcgcttcc cgcttcctcg 5940
 E--> 603 ctcactgact cgctcgctc ggtcggttccg ctgcggcgg 5940
 E--> 605 gcggtataac gtttatccac agaattcagg gataacgcag gaaagaaatc gtgagcaaaa 6060
 E--> 607 ggccagcaaa aggccaggaa ccgtaaaaag gcccgttgc tggcggtttt ccattggctc 6120
 E--> 609 cgccccccctg acgagcatca caaaaatcga cgctcaagtc agagggtggcg aaacccgaca 6180
 E--> 611 ggactataaa gataccaggc gtttccctt ggaagctccc tcgtcgctc tcctgttccg 6240
 E--> 613 accctgccc ttaccggata cctgtccggc tttccctt cgggaaggctt ggcgtttct 6300
 E--> 615 catagctcac gctgttaggtat ttcgttccg gtgttaggtc ttcgttccaa gctgggctgt 6360
 E--> 617 gtgcacgaaac ccccggttca gcccggaccgc tgcccttat ccggtaacta tcgttccg 6420
 E--> 619 tccaaacccgg taagacacga cttatcgca ctggcggcag ccactggtaa caggatttgc 6480
 E--> 621 Page 9
 W--> 625 70094 Syngenta *delete*
 E--> 627 agagcgaggt atgttagggcgg tgctacagag ttcttgaagt ggtggctaa ctacggctac 6540
 E--> 629 actagaagaa cagtatttgg tatctgcgt ctgtgtttt cggaaaaaaga 6600
 E--> 631 gttggtagct cttgtatccgg caaacaacc accgctggta gccgtgggtt tttttttgc 6660
 E--> 633 aagcagcaga ttacgcgcag aaaaaaagga tctcaagaag atcctttgtat ctttctacg 6720
 E--> 635 gggctgtacg ctcaatggaa cgaaaactca cggtttaaggaa ttttggatcat gagattatca 6780
 E--> 637 aaaaggatct tcacccatgt cttttgtatc cggtttaat tccctgtgggtt ggcattgcaca 6840
 E--> 639 tacaatggaa cgaacggata aaccccccata cggcccttta aatatccat tattctaata 6900
 E--> 641 aacgctctt tctcttaggt ttacccggca atatatccgt tcaaaacactg atagtttaaa 6960
 E--> 643 ctgaaggcgg gaaacgacaa tctgtatcatg agcggagaat taaggagtc acgtttagac 7020
 E--> 645 ccccgccgt gacgcgggac aagccgtttt acgtttggaa ctgacagaac cgcaacgcgt 7080
 E--> 647 caggaatttgg cccgtatccggc cattttaaatc aattttggcgc gtacgtatca ctatgtcg 7140
 E--> 649 atcgcttaat taagccggcgc gcctaaagct tctggcagac aaagtggcag acataactgtc 7200
 E--> 651 ccacaaatga agatgaaatc tgtaaaaagaa aacgcgtgaa ataatgcgtc tgacaaagg 7260
 E--> 653 taggtcggtc gcctttaatc aataccaaaatg tggccctac cacgtatggaa aaactgtgca 7320
 E--> 655 gtcgggttgg cttttcttgc cgaacaaata agatcggtt ccgacaggtg ggggtccacc 7380
 E--> 657 atgtgaaggc atcttccatc tccaaataatg gagcaatgac gtaagggtt acgaaataag 7440
 E--> 659 taagggtat ttggaaaatg tccactcacc cgtcaatgtata taaatactt gcccctccct 7500
 E--> 661 cattgttaag ggagcaagga tccaccggc gcccattatgg ccctgtccaa caagtttcatc 7560
 E--> 663 ggccacgaca tgaagatgac ctaccacatg gacggctgcg tgaacggcca ctacttcacc 7620
 E--> 665 gtgaaggcgg agggcggccgg caagccctac gagggcacc acacccatc cttcaagg 7680
 E--> 667 accatggcca acggccggcc cctggccctc tccttcgaca tcctgtccac cgttgcgt 7740
 E--> 669 tacggcaacc gctgttccac cgccttccccc accagcatgc ccgactactt caagcaggcc 7800
 E--> 671 ttccccgacg gcatgttccatc cgagagaacc ttccatcagc aggacggcgg cgtggccacc 7860
 E--> 673 gccagctggg agatcagccct gaaggggcaac tgcttcgac acaagtccac cttccacggc 7920
 E--> 675 gtgaacttcc cccggacgg cccctgtatc gccaagaaga ccacccggctg ggacccctcc 7980

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/517,039

DATE: 12/22/2004
TIME: 09:15:16

Input Set : A:\Sequence Listing-70094USPCT.txt
Output Set: N:\CRF4\12222004\J517039.raw

E--> 677 ttcgagaaga tgaccgtgtc cgacggcatc ttgaaggcg acgtgaccgc cttcctgatg 8040
 E--> 679 ctgcagggcg gcgcaacta cagatccag ttccacaccc cctacaagac caagaagccc 8100
 E--> 681 gtgaccatgc ccccaacca cgtgggtggag caccgcac ccagaacccga cctggacaag 8160
 E--> 683 ggcggcaaca gcgtgcagct gaccgagcac gccgtggccc acatcaccc cgtggtgccc 8220
 E--> 685 ttctgagagc tctagatccc cgaatttccc cgatcgatca aacatttgc aataaagttt 8280
 E--> 687 cttaaagattt aatccgttg ccggcttgc gatgattatc atataatttc tggatgat 8340
 E--> 689 cgtaaagcat gtaataattt acatgtatg catgacgtt tttatgatg gggttttat 8400
 E--> 691 Page 10
 W--> 694 70094 Syngenta delete

SAME
ERORS

E--> 696 gattagagtc cgcattat acatttata cgcgatagaa aacaaaatat agcgcgcaaa 8460
 E--> 698 ctaggataaa ttatcgccg cgggttcata tatgttacta gatcggaat tgggtaccga 8520
 E--> 700 attcaactggc cgtcggttta caacgtcgtg actggaaaa ccctggcggtt acccaactta 8580
 E--> 702 atcgcccttc agcacatccc ccttcgcga gctggcgtaa tagcgaagag gccgcacccg 8640
 E--> 704 atcgcccttc ccaacagttt cgcagcctga atggcgaatg ggcgcgtatg cggatatttc 8700
 E--> 706 tccttacgca tctgtcggtt atttcacacc gcataatggtg cactctcagt acaatctgt 8760
 E--> 708 ctgatgccgc atagtaagc cagccccgac acccgccaa acccgctgac gcccgcgtac 8820
 E--> 710 gggctgtct gctccggca tccgcttaca gacaagctgt gaccgtctcc gggagctgca 8880
 E--> 712 tgggtcgag gtttccaccg tcatcaccga aacgcgcgag acgaaaggc ctcgtgatac 8940
 E--> 714 gcctatttt ataggttaat gtcgtatataa taatggttt ttagacgtca ggtggcactt 9000
 E--> 716 ttcggggaaa tgtgcgcggaa accccctattt gtttattttt ctaaatatcat tcaaatatgt 9060
 E--> 718 atccgctcat gagacaataa ccctgataaa tgcttcaatg ggcgcgggtt accagcttgc 9120
 E--> 720 atgcctgcag gtcgactcta gaggatcccg gcagacaaag tggcagacat actgtcccac 9180
 E--> 722 aaatgaagat ggaatctgtt aagaaaaacg cgtgaaataa tgcgtctgac aaaggtagg 9240
 E--> 724 tcggctgcct ttaatcaata ccaaagtgtt ccctaccacg atggaaaaac tgtgcagtcg 9300
 E--> 726 gtttggcttt ttctgacgaa caaataagat tgcgtggccga caggtggggg tccaccatgt 9360
 E--> 728 gaaggcatct tcagactcca ataatggagc aatgacgtaa gggcttacga aataagtaag 9420
 E--> 730 ggttagttgg gaaatgtcca ctcacccgtc agtctataaa tacttagccc ctccctcatt 9480
 E--> 732 gttaagggag caaaatctca gagagatgt cctagagaga gaaagagagc aagtagccta 9540
 E--> 734 gaagta 9546

737 <210> SEQ ID NO: 9543
 738 <211> LENGTH: 10604
 739 <212> TYPE: DNA
 740 <213> ORGANISM: Artificial
 742 <220> FEATURE:
 743 <223> OTHER INFORMATION: pBSC11234
 745 <400> SEQUENCE: 3

746 cgccctaa gcttgcatgc ctgcaggcg actctagagg atccctggcag acaaagtggc 60
 748 agacatactg tcccacaaat gaagatggaa tctgtaaaag aaaacgcgtg aaataatgcg 120
 750 tctgacaaag gtttagtgcgg ctgcctttaa tcaataccaa agtggtcctt accacgatgg 180
 752 aaaaactgtt cagtcgggtt ggcttttctt gacgacaaa taagattgt ggccgacagg 240
 754 tgggggtcca ccatgtgaag gcatcttgcg actccaataa tggagcaatg acgtaaaggc 300
 756 ttacgaaata agtaagggta gtttggaaa tgcctactca cccgtcagtc tataaataact 360

E--> 758 Page 11
 W--> 760 70094 Syngenta delete

E--> 762 tagccctcc ctcattgtta agggagccaa atctcagaga gatagtccta gagagagaaa 420
 E--> 764 gagagcaagt agcctagaag taggatcccc gatcatgca aaactcatca actcagtcgca 480
 E--> 766 aaactatgcc tggggcagca aaacggcggtt gactgaactt tatggatgg aaaatccgtc 540
 E--> 768 cagccagccg atggccgagc tggatggg cgacatccg aaaagcaattt caccagtgca 600
 E--> 770 gaatgccgccc ggagatatcg tttcactgcg tgcgtgtt gggatgtata aatcgactct 660

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/517,039

DATE: 12/22/2004
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Input Set : A:\Sequence Listing-70094USPCT.txt
Output Set: N:\CRF4\12222004\J517039.raw

E--> 772 gctcgagag gccgttgc aacgcttgg cgaactgcct ttcctgttca aagtattatg 720
 E--> 774 cgcagcacag ccactctcca ttcaggttca tccaaacaaa cacaattctg aatcggttt 780
 E--> 776 tgccaaagaa aatgcgcag gtatcccgtt ggatggcc gaggtaat ataaagatcc 840
 E--> 778 taaccacaag ccggagctgg ttttgcgtt gacgccttc ctgcgtatga acgcgttgc 900
 E--> 780 tgaattttcc gagattgtct ccctactcca gccggtegaa ggtgcacatc cggcgattgc 960
 E--> 782 tcactttta caacagcctg atgccaacg ttaagcgaa ctgttcgcca gcctgttcaa 1020
 E--> 784 tatgcagggt gaagaaaaat cccgcgtt ggcgatttttta aatcgcccc tcatagccca 1080
 E--> 786 gcagggtgaa ccgtggaaaa cgttgcgtt aatttctgaa ttttaccgg aagacagcgg 1140
 E--> 788 tctgttctcc ccgctattgc tgaatgtgtt gaaattgaac cctggcgaag cgatgttcc 1200
 E--> 790 gttcgctgaa acacccacg cttacctgca aggctgtggcg ctggaaatgaa tggaaactc 1260
 E--> 792 cgataacgtg c69d9gcgg gtctgacgcc taaatacatt gatattccgg aactgggtgc 1320 1318
 E--> 794 caatgtgaaa ttcgaagcca aaccggctaa ccagttgtt acccagccgg tgaacaagg 1380
 E--> 796 tgcagaactg gacttccgaa ttccagtggaa tgattttgc ttctcgatgc atgaccttag 1440
 E--> 798 tgataaaagaa accaccatta gccagcagag tgccgcatt ttgttctgcg tgaaggcga 1500
 E--> 800 tgcaacgttggaaagggtt ctcagcaggta acagcttaaa cgggtgaat cagcgtttat 1560
 E--> 802 tgccgccaac gaatcaccgg tgactgtcaa aggccacggc cggttagcgc gtgtttacaa 1620
 E--> 804 caagctgtaa gagcttactg aaaaaattaa catctttgc taagctggaa gctctagatc 1680
 E--> 806 cccgaatttc cccgatcggtt caaacatttgc gcaataaagt ttcttaagat tgaatccgt 1740
 E--> 808 tgccggctt gcgatgatta tcatataatt tctgttgaat tacgttaagc atgatataat 1800
 E--> 810 taacatgtaa tgcgtacgt tattttatgag atgggtttt atgatttagag tcccgcaatt 1860
 E--> 812 atacatttaa tacgcgtatg aaaacaaaat atagcgcgc aactaggata aattatcgcg 1920
 E--> 814 cgggtgtca tctatgttac tagatggga attgggtacc atgcccggc ggcagcatg 1980
 E--> 816 gccgtatccg caatgtgttta ttaagttgtc taagcgtaa ttgttttaca ccacaatata 2040
 E--> 818 tcctgccacc agccagccaa cagctcccg accggcagct cggcacaaaa tcaccactcg 2100
 E--> 820 atacaggcag cccatcagaa ttaattctca ttttgcgtt gtttgcgtt gtttgcgtt 2160
 E--> 822 gcaccaatgc ttctggcgcc aggcagccat cgaaagctgt ggtatggctg tgcaggctg 2220
 E--> 824 aaatcaatgc taaattcgtg tgcgtcaagg #gcactcccg ttctggataa tttttttgc 2280

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delete

E--> 826
 W--> 828
 E--> 830 gcccacatca taacggttct ggcaaatatt ctgaaatgag ctgttgcacaa ttaatcatcc 2340
 E--> 832 ggctcgata atgttgaa ttgtgagccg ataacaattt cacacaggaa acagaccatg 2400
 E--> 834 agggaaagct tgatcgccga agtatcgact caactatcg aggtagttgg cgtcatcgag 2460
 E--> 836 cggccatctcg aaccgacgtt gctggccgtt cattgtac gctccgcagt ggatggccgc 2520
 E--> 838 ctgaagccac acagtgtat tgatgttgcgtt gttacggta ccttccctgg tgatgaaaca 2580
 E--> 840 aecggcgcag ctttgcataa cgaccttttgc gaaacttcgg agagagcgg 2640
 E--> 842 attctccgcg ctgtagaagt caccattgtt gtgcacgc acatcatcc gtggcgat 2700
 E--> 844 ccagctaagc gcgaaactgca atttggagaa tggcagcgc atgacatctc tgcaaggatc 2760
 E--> 846 ttgcgagccag ccacgatcga cattgtatcg gctatcttc tgacaaaagc aagagaacat 2820
 E--> 848 agcgttgcct tggttaggtcc agcggcggag gaactcttgc atccggatcc tgaacaggat 2880
 E--> 850 ctatggagg cgctaaatga aaccttaacg ctatggact cggcccccga ctggctggc 2940
 E--> 852 gatgagcga atgttagtgcg ttcgttgc cgcattttgtt acagcgcagt aaccggcaaa 3000
 E--> 854 atcgcccgaa aggtatgcgc tgccgactgg gcaatggagc gcctgcccggc ccagtatcag 3060
 E--> 856 cccgtcatac ttgaagcttag gcaggctt cttggacaag aagatcgatc ggctcgccgc 3120
 E--> 858 gcagatcgt tggaagaatt ttttgcgtt gtttgcgtt agatcaccaa agtagcggc 3180
 E--> 860 aaataaaagct ctagtggatc tccgttcccg cggggatct ggctcgccggc ggacgcacga 3240
 E--> 862 cgccggggcg agaccatagg cgatctccca aatcaatagt agctgttacc tgcaggctt 3300
 E--> 864 tcactgtttaa caacgattga gaattttgtt ctttttttttgcgtt aaatacttgcgtt 3360
 E--> 866 tgcgtatccgc ggtcagccgc aattctgcgtt aactgcgtt ttagctggag atgattgtac 3420
 E--> 868 atcccttcacg tgaaaatttc tcaagcgtt gtttgcgtt agattgaaag 3480

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/517,039

DATE: 12/22/2004
TIME: 09:15:16

Input Set : A:\Sequence Listing-70094USPCT.txt
Output Set: N:\CRF4\12222004\J517039.raw

INVALID
Amino
Acid
designators

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/517,039

DATE: 12/22/2004
TIME: 09:15:16

Input Set : A:\Sequence Listing-70094USPCT.txt
Output Set: N:\CRF4\12222004\J517039.raw

E--> 968 tcctggtata ggtctgcata tccgactcgt ccaacatcaa tacaacctat taatttcccc 6240
E--> 970 tcgtcaaaaa taagggttatac aagtggaaaaa tcaccatgag tgacgactga atccggtag 6300
E--> 972 aatggcaaaa gctctgcatt aatgaatcgg ccaacgcgcg gggagaggcg gtttgcgtat 6360
E--> 974 tggcgctct tccgcttcct cgctcaactga ctcgcgcgc tcggcgttc ggctgcggcg 6420
E--> 976 agcggtatca gctcaactcaa aggccgtaat acggttatcc acagaatcag gggataacgc 6480
E--> 978 aggaaagaac atgtgagcaa aaggccagca aaaggccagg aaccgtaaaa aggccgcgtt 6540
E--> 980 gctggcggtt ttccataggc tccgcccccc tgacgagcat cacaatc gacgctcaag 6600
E--> 982 tcagagggtgg cgaaacccga caggactata aagataccag gcgtttcccc ctggaaagctc 6660
E--> 984 cctcgtgcgc tctccgttcc cgaccctgcc gcattaccgg tacctgtccg cctttctccc 6720
E--> 986 ttcgggaagc gtggcgcttt ctcatagctc acgctgttagg tatctcagtt cggtaggt 6780
E--> 988 cgttcgtcc aagctgggt gtgtgcacga accccccgtt cagccccgacc gctgcgcctt 6840
E--> 990 atccggtaac ttcgtcttg agtccaaaccc ggtaagacac gacttacgc cactggcagc 6900
E--> 992 agccactggt aacaggatta gcagagcggag gtatgttagc ggtgctacag agttcttgaa 6960
E--> 994 gtggggccct aactacggct acactagaag aacagtattt ggtatctgcg ctctgctgaa 7020
E--> 996 gccagttacc ttccggaaaaa gagttggtag ctcttgatcc ggcaaaacaaa ccaccgctgg 7080
E--> 998 tagcggtggg tttttgttt gcaagcagca gattacgcgc agaaaaaaag gatctcaaga 7140
E--> 1000 agatcctttg atctttctta cggggctcga cgctcagtgg aacgaaaact cacgttaagg 7200
E--> 1002 gattttggc atgagattat caaaaaggat cttcacctag atcctttga tccggaaatta 7260
E--> 1004 attcctgtgg ttggcatca catacaaatg gacaacggg taaacctttt cacgcctttt 7320
E--> 1006 taaatatccg attattctaa taaacgtct ttttctttag gtttaccgc caatatatcc 7380
E--> 1008 tgtcaaacac tgatagttt aactgaaggc gggaaacgc aatctgatca tgagcggaga 7440
E--> 1010 attaaggag tcacgtttag accccccgg atgacgcggg acaagccgtt ttacgtttg 7500
E--> 1012 aactgacaga accgcaacgc tgcaggatt ggcgcgcgc gccattaaa tcaattggc 7560
E--> 1014 gcgtacgtag cactgtgcg cgatcgctt attaagcggc ggcgcctgcag gcccgcac 7620
E--> 1016 aattattata tcaaaatggc aaaaacattt aatacgtatt atttaagaaa aaaatatgt 7680
E--> 1018 ataatatatt tatattttaa tatctattct tatgtatttt taaaaatct attatatatt 7740
E--> 1020 gatcaactaa aatatttttata ttttacact ttttgcatttttataat ttttgcgt 7800
E--> 1022 ttttggcat attaataat gactattttttaataatcga tcattattct tacatggtac 7860
E--> 1024 atattgtgg aaccatata gatgtccatt gatgtgatc atgtggatag tttttgtac 7920
E--> 1026 Page 15
W--> 1029 70094 Syngenta *delete*
E--> 1031 caggcctcca tttggcgctt attaattaaat ttgttaacag tccgtactaa tcagttactt 7980
E--> 1033 atccttcctc catataatt aatcttggta gtctcgaatg ccacaacact gactagtctc 8040
E--> 1035 ttggatcata agaaaaagcc aaggaacaaa agaagacaaa acacaatggg agtacccctt 8100
E--> 1037 gcatagcaat gtctaaatgc ataaaatca aacaaaaacg caatcacaca cagtggacat 8160
E--> 1039 cacttatcca ctatgtgatc aggtatcggc cgtcaagaaa aaaaactgg accccaaaag 8220
E--> 1041 ccatgcacaa caacacgtac tcacaaaagggt gtcacatcgag cagccaaaa cattcaccaa 8280
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E--> 1045 cttccgcac ctcatttttgc ttatattca caccgcgtca actgcacatc accccgtggc 8400
E--> 1047 caaatgtcca tgcatttttgc caagacccat gactataat atctgcacatc tcggccagg 8460
E--> 1049 ttttcatcat caagaaccag ttcaatattcc tagtacaccg tattaaagaa ttaagatat 8520
E--> 1051 actccacccgg atccaccatg gccaagctag tttttccct ttgtttctg ctttcagtg 8580
E--> 1053 gctgtcgctt cgctggat ttcggcaaga cttccgcga gggccgcctt gtgctcaagg 8640
E--> 1055 agaagaactt caccgtggag ttccggcggtt agaagatcca cctcggctgg aagatatcg 8700
E--> 1057 gccgcgtgaa gggctcgccg ggccgcctcg aggtgcgtccg caccaaggcc cccggagaagg 8760
E--> 1059 tgctcgtaa caactggcag ttccggggcc cgtccgcgtt ggtggacgccc ttctccttca 8820
E--> 1061 agccgcgcga gatgcacccg aactggcgct acaccgcac cgtggcgccg gacgtgcgt 8880
E--> 1063 agcgcacact gcaatggcgtt tacttcgtgg ccgaggaggg caaggtgtac ggttccctt 8940
E--> 1065 cctccaagat cgcgcgcgc ttctcgcgg tggaggacgg cgagctggcgc 9000

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Output Set: N:\CRF4\12222004\J517039.raw

E--> 1067 agtacttcga cgtggagttc gacgacttcg tgccgcttgg a g c c g c t c g t g g t g c t c g a g g 9060
E--> 1069 acccgAACAC cccgtccctc ctcgagaagt acggcagact ggtgggcatg gagaacaacg 9120
E--> 1071 cccgggtgcc gaagcacacg ccgaccggct ggtgctcctg gtatcaactac ttccctcgacc 9180
E--> 1073 tcacctggga ggagaccctc aagaacctca agctcgccaa gaacttcccg ttcgaggtgt 9240
E--> 1075 tccagatcga cgacgcctac gagaaggaca tcggcactg gctcgtgacc cgcggcgact 9300
E--> 1077 tcccgtccgt ggaggagatg gccaagggtga tcggccgagaa cggcttcata cccggcatct 9360
E--> 1079 ggaccgcccc gtttcctcg tccgagacta gtgacgtgtt caacgagcac cccggactggg 9420
E--> 1081 tggtaagga gaacggcgag ccgaagatgg cctaccgc aa ctggaaacaag aagatttacg 9480
E--> 1083 ccctcgacct ctccaaggac gaggtgctca actggctt c g a c c t t c t c t c t c 9540
E--> 1085 gcaagatggg ctaccgc t a c t c a g a t c g a c t t c t t c t c t c t c 9600
E--> 1087 agcgcaagaa gaacatcacc ccgatccagg cctccgc a a g g c a t c g a g a c c a t c c g c a 9660
E--> 1089 aggccgtggg ggaggactcc ttcatcctcg gtcggcgc c c c c t t c t c t c c g g c g t g g 9720
E--> 1091 gtcgcttgg a t g g c a t g c g c a t c g g c c c g g a c a c a t c g 9780
E--> 1093 aggacaacgg egcccccggc gcccgttggg ccctccgc a a c c a t c a c c c g t a c t t c a 9840

Page 16

70094 Syngenta

del/ete

E--> 1095
W--> 1098
E--> 1100 tgcacgaccg cttctggctc aacgaccgg actgcctcat cctccgcag gagaagaccg 9900
E--> 1102 acctcacca gaaggagaag gagctgtact cctacacctg cggcgatcta gacaacatga 9960
E--> 1104 tcatcgagtc cgacgacctc tccctcg tgcgaccacgg caagaagggtg c t c a a g g a g a 10020
E--> 1106 ccctcgagct gtcggggc aggccgcg tgcagaacat catgtccgag gacctccgct 10080
E--> 1108 acgagatcgt gtcctcg g accctctcg gcaacgtgaa gatcgtggg gacctaact 10140
E--> 1110 cccgcgagta ccaccccg aaggaggca agtccctccct caagaagcgc g t g g t g a a g c 10200
E--> 1112 gggaggacgg caggaacttc tacttctacg aggaggcga g c g c g a g t g a a g c t t g a c g 10260
E--> 1114 tcaactagtgc gatcgcgct a g c c a t g g c g c g c c c g g a g a t c c c g a a t t c 10320
E--> 1116 cccgatcggtt caaaatattt gcaataaagt ttcttaagat t g a a t c t g t t g c c g g t c t t 10380
E--> 1118 gcatgattataatt tctgttgaat tacgtttagc atgtaataat taacatgtaa 10440
E--> 1120 tgcattgttgcgatg tattttatgat atgggtttt atgatttagag t c c c g a a t t a t a a t 10500
E--> 1122 tacgcgatag aaaacaaaat atagcgccca aactaggata aattatcg g c g g g t g t c a 10560
E--> 1124 tctatgttac tagatcg g a t t c c t c g a g t c t a g a c t g t 10604

1127 <210> SEQ ID NO: 4
1128 <211> LENGTH: 8755
1129 <212> TYPE: DNA

Found 8755

10602

1130 <213> ORGANISM: Artificial
1132 <220> FEATURE:
1133 <223> OTHER INFORMATION: pBScl1369

1135 <400> SEQUENCE: 4

1136 aagcttctgg cagacaaagt ggcagacata ctgtcccaca aatgaagatg gaatctgtaa 60
1138 aagaaaacgc gtgaaataat g c g t c t g a c a a g g t t a g g t c g g c t c t t a a t c a a t a c 120
1140 caaagtggtc cctaccacga tggaaaact gtgcagtcgg t t t g g c t t t t t c t g a c g a a c 180
1142 aaataagatt cgtggccgac aggtgggggt ccaccatgtg a a g g c a t c t c a g a c t c c a a 240
1144 taatggagca atgacgtaag ggcttaccaa a t a a g t a a g g g t a g t t t g g g a a t g t c c a c 300
1146 tcaccctgtca gtctataat acttagcccc tccctcattt g t t a a g g g a g c a a g g a t c c a c 360
1148 cggtcgcccc catggccca g t c c a a g c a c g g c t g a c c a a g g a t g a a g t a c c 420
1150 gcatggaggg ctgcgtggac ggccacaagt tcgtgatcac c g g c g a g g g c a t c g g c t a c c 480
1152 ctttcaaggg caagcaggcc atcaacctgt g c g t g g t g g a g g c g g c c c c t t c g 540
1154 ccgaggacat cttgtccgccc g c c t t c a t g t a c g g c a a c c g c g t g t t c a c c 600
1156 aggacatcgt cgactacttc a a g a a c t c t c g c c c g g c t a c a c t t g g a c c g c t c c t 660
1158 tccctgttca ggacggcgcc g t g t g c a t t g a c a c g c c g a c a t c a c c g t g a g c g t g g a g g 720
1160 agaactgcattt g t a c c a c g a g t c a a g t t c t a c g g c t c c t a c c c c g c c 780

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Output Set: N:\CRF4\12222004\J517039.raw

E--> 1162 Page 17
W--> 1164 70094 Syngenta *del file*
E--> 1166 ttagaagaa gatgaccgac aactgggagc cctcctgcga gaagatcatc cccgtgccc 840
E--> 1168 agcagggcat cttgaaggc gacgtgagca tgtacctgct gctgaaggac ggtggccgct 900
E--> 1170 tgcgctgcga gttcgacacc gtgtacaagg ccaagtccgt gcccccaag atgcccgact 960
E--> 1172 ggcacttcat ccagcacaag ctgaccccgag aggaccgcag cgacgccaag aaccagaagt 1020
E--> 1174 ggcacctgac cgagcacgac atcgccctcg gctccgcctt gccctgtct agatcccga 1080
E--> 1176 tttccccgtat cgttcaaaca tttggcaata aagtttctta agattgaatc ctgttgccgg 1140
E--> 1178 tcttgcgtat attatcatat aatttctgtt gaattacgtt aagcatgtaa taattaacat 1200
E--> 1180 gtaatgcgt acgttattta tgagatgggt tttatgatt agagtcccgc aattatacat 1260
E--> 1182 ttaatacgcg atagaaaaca aaatatacg cgccaaactag gataaattat cgccgcgg 1320
E--> 1184 gtcatctatg ttactagatc gggaaatttggg gaaatttacc ggtgccaat tttccgcata 1380
E--> 1186 cagcttctgg cagacaaagt ggcagacata ctgtcccaca aatgaagatg gaatctgtaa 1440
E--> 1188 aagaaaacgc gtgaaaataat gctgtgaca aaggttaggt cggctgcctt taatcaatac 1500
E--> 1190 caaagtggtc cctaccacga tggaaaaact gtgcagtcgg tttggcttt tctgacgaac 1560
E--> 1192 aaataagatt cgtggccgac aggtgggggt ccaccatgtg aaggcatctt cagactccaa 1620
E--> 1194 taatggagca atgacgtaa ggcttacgaa ataagtaagg gtatggggg aatgtccac 1680
E--> 1196 tcaccctgtca gtctataat aattagcccc tccctcattt ttaaggagc aaggatccat 1740
E--> 1198 gaaaaagct gaactcaccg cgacgtctgt cgagaagttt ctgatgaaa agttcgacag 1800
E--> 1200 cgtctccgac ctgatgcgc tctcgaggg cgaagaatct cgtcttca gcttcgatgt 1860
E--> 1202 aggagggcgt ggatatgtcc tgccggtaaa tagctgcgc gatggttct acaaagatcg 1920
E--> 1204 ttatgtttat cggcaacttgc catcgccgc gctcccgatt ccggaaagtgc ttgacattgg 1980
E--> 1206 ggaatttcgcg gagagcctga cctattgcattt ctcccgccgt gcacagggtg tcacgttgc 2040
E--> 1208 agacctgcct gaaaccgaaac tgcccgtgt tctcgagccg gtcgcggagg ccatggatgc 2100
E--> 1210 gatcgctgcg gccgatcttta gccagacgag cgggttcggc ccattcggac cgcaaggaat 2160
E--> 1212 cggtaatac actacatggc gtgatttcattt atgcgcgatt gctgatcccc atgtgtatca 2220
E--> 1214 ctggcaaact gtgatggacg acaccgtcag tgcgtccgtc ggcgcggctc tcgatgagct 2280
E--> 1216 gatgctttgg gccgaggact gccccgaagt cggcacctc gtgcacgcgg atttcggctc 2340
E--> 1218 caacaatgtc ctgacggaca atggccgcat aacagcggtc attgactgga gcgaggcgat 2400
E--> 1220 gttcgggat tcccaatacg aggtgcacaa catcttc tggaggccgt ggttggctt 2460
E--> 1222 tatggagcag cagacgcgtc acttcgacgc gaggcatccg gagcttcgag gatcggccgc 2520
E--> 1224 gctccggcgt tatatgtcc gcattggctt tgaccaactc tatcagagct tggttgcgg 2580
E--> 1226 caatttcgtat gatgcagctt gggcgcaggc tgatgcgc gcaatgtcc gatccggagc 2640
E--> 1228 cgggactgtc gggcgatcac aaatcgcccg cagaagcgac gccgtctgga ccgatggctg 2700
E--> 1230 Page 18
W--> 1233 70094 Syngenta *del file*
E--> 1235 ttagaagta ctcggccata gtggaaaccc acggcccgac actcgccga *99* *2760* *2758 this?*
E--> 1237 atagggtatcc cccgaatttc cccgatcgat ctttgcgtt caaacatttgc gcaataaaatg ttttgcgtt 2820
E--> 1239 tgaatcctgt tgccggctt gcgatgatataatt ttttgcgtt tacgttgcgc 2880
E--> 1241 atgtaataat taacatgtaa tgcatgcgtt tatttgcgtt atgggtttt atgatttagag 2940
E--> 1243 tcccgcaatt atacatttaa tacgcgtatc aaaaacaaat atgcgcgc aactaggata 3000
E--> 1245 aattatcgac cgcgggtgtca ttttgcgtt tagatcgccg attagccgc cgaatttact 3060
E--> 1247 ggcgcgtt ttacaacgtc tgactggga aaaccctggc gttaccaac ttaatgcgc 3120
E--> 1249 tgcacgcacat cccctttcg ccaggccgg ccacgttgcgc cgtatccgc atgtgttatt 3180
E--> 1251 aagttgttca agcgtcaatt tggttgcgtt acaatatacg ctgcacccag ccagccaaaca 3240
E--> 1253 gctccggcgt cggcagctcg gcacaaaatc accactcgat acaggcagcc catcagaatt 3300
E--> 1255 aatttcgtat tttgcgtt tacgttgcgc accaatgtt ctggcgatcg 3360
E--> 1257 gcaaggccatcg gaagctgtgg tatggctgtg caggtcgtaa atcactgtat aattcgtgtc 3420
E--> 1259 gctcaaggcg cactcccgat ttggataatg tttttgcgc cgacatcata acgggttctgg 3480

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Output Set: N:\CRF4\12222004\J517039.raw

E--> 1261 caaatattct gaaatgagct gttgacaatt aatcatccgg ctcgtataat gtgtggaatt 3540
 E--> 1263 gtgagcggat aacaatttca cacagggaaac agaccatgag ggaagcggtt atcgccgaag 3600
 E--> 1265 tatcgactca actatcagag gtagttggcg tcatcgagcg ccatctcgaa ccgacggttgc 3660
 E--> 1267 tggccgtaca tttgtacggc tccgcagttt atggcggtt gaagccacac agtgatattt 3720
 E--> 1269 atttgcgtt tacgttgacc gtaaggctt atgaaacaac gggcgagct ttgatcaacg 3780
 E--> 1271 accttttggg aacttcggct tcccctggag agagcgagat tctccgcgt gttagaagtca 3840
 E--> 1273 ccattgttgt gcacgacgac atcattccgt ggcgttatcc agctaagcgc gaactgcaat 3900
 E--> 1275 ttggagaatg gcagcgcaat gacattctt caggtatctt cgagccagcc acgatcgaca 3960
 E--> 1277 ttgatctggc tatcttgcgt aaaaaagcaa gagaacatag cggtgcctt gttagtccag 4020
 E--> 1279 cggcggagga actctttgtt ccggttcctt aacaggatct atttgaggcg ctaaatgaaa 4080
 E--> 1281 ccttaacgct atggaactcg ccgcccgaact gggctggcga tgagcgaat gtgtgccta 4140
 E--> 1283 cgttgcggc cattgggtac agcgcaatcc cggccaaat cgcgcgaag gatgtcgctg 4200
 E--> 1285 ccgactgggc aatggagcgc ctgcggggcc agtatacgcc cgtcataactt gaagcttaggc 4260
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 E--> 1291 cgtacccggg gatctggctc gggcgccgacg cacgacgccc gggcgagacc ataggcgatc 4440
 E--> 1293 tcctaaatca atatgtactt taacctcgaa gcgttactt tctaacaacg attgagaattt 4500
 E--> 1295 tttgtatctt aatggaaata cttggttcgc attttgtca tccgcgtca gccgcaattt 4560
 E--> 1297 Page 19
 W--> 1300 70094 Syngenta *del/te*

E--> 1302 tgacgaactg cccatattgc tggagatgtatgtatcct tcacgtaaaa atttctcaag 4620
 E--> 1304 cgctgtgaac aagggttgcg attttagatt gaaaggttag cgcgttgcgaa acgttcttct 4680
 E--> 1306 tgcgtatgac gacgtcgctt tgcggcatct tattattgaa taccttacga tccacgcctt 4740
 E--> 1308 caaagtgacc gcggttagccg acagcacccca gttcacaaga gtactctt ccgcgcacgg 4800
 E--> 1310 cgatgtcggt gttgttgcgt tagatttagg tcgtgaagat gggctcgaga tcgttcgtaa 4860
 E--> 1312 tctggccggca aagtctgata ttccaatcat aattatcagt ggcgaccggc ttgaggagac 4920
 E--> 1314 ggataaaagt gttgcactcg agctaggagc aagtgtttt atcgctaaacg cgttcgttat 4980
 E--> 1316 cagagagttt ctgcacgcgat ttcgggttgc cttgcgcgtg cggcccaacg ttgtccgcctc 5040
 E--> 1318 caaagacccga cggctttttt gtttacttgc ctggacactt aatctcaggc aacgtcgctt 5100
 E--> 1320 gatgtccgaa gctggccggc aggtgaaact tacggcaggt gaggtaatc ttctccgc 5160
 E--> 1322 gtttttagag aaaccccgccg acgttctatc gcgcgagccaa cttctcattt ccagtcgagt 5220
 E--> 1324 acgcgcacgag gaggtttatg acaggagttt agatgttctc attttgaggc tgcgcgc 5280
 E--> 1326 acttgaggca gatccgtcaa gcccctcaact gataaaaaaca gcaagaggtt cccgttattt 5340
 E--> 1328 ctttgacgcg gacgtgcagg tttgcacgg ggggacgtat gcagccttag ccaattccca 5400
 E--> 1330 gatccccggag gaatcggttgc gaggcggttgc aaaccatccg gcccgttaca aatcgccgc 5460
 E--> 1332 gcgctgggttgc atgacctgttgc gggaaagttt aaggccgcgc aggcccccac gggcaacgc 5520
 E--> 1334 atcgaggcag aagcacgcggc cggtaatcg tggcaagcgcc cgcgtatcg aatccgc 5580
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 E--> 1346 taccggggaa ggaaggggaga caagccccgc cggcggttcc gtccacacgt tgcggacgt 5940
 E--> 1348 ctcaagtttgc gccggcgacg cgtatggcgga aaggcagaaag acgacctgttgc agaaacactgc 6000
 E--> 1350 attcggtttaa acaccacgcg cgttgcctatc cagcgatcgaa agaaggccaa gaacggccgc 6060
 E--> 1352 ctgggtacgg tatccgaggg tgaaggcttgc attagccgtt acaagatcgta aaagagcgaa 6120
 E--> 1354 accggccggc cggagatcatcg cggatcgatc ctagctgttgc ggtatgtaccg cgagatcaca 6180
 E--> 1356 gaaggcaaga accccggacgt gctgacgttgc caccggattt acttttgcgttgc gatccggc 6240
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Output Set: N:\CRF4\12222004\J517039.raw

E--> 1360 ttgttcaaga cgatctacga acgcagttgc aggcggag agttcaagaa gttctgtttc 6360
 E--> 1362 accgtgcgca agctgatcgg gtcaaatgac ctggcggagt acgatttcaa ggaggaggcg 6420
 E--> 1364 gggcggctg gcccgtt agtcatgcgc/taccccaacc tgatcgaggg cgaagcatcc 6480
 E--> 1366 Page 20
 W--> 1368 70094 Syngenta *delete*
 E--> 1370 gccggttcct aatgtacgga gcagatgcta gggcaaattt ccctagcagg gaaaaaaagg 6540
 E--> 1372 cggaaaaggc tctttcctgt ggatagcag tacattggg acccaaagcc gtacattggg 6600
 E--> 1374 aaccggaaacc cgtacattgg gaaacccaaag cgtacattt ggaacccggc acacatgtaa 6660
 E--> 1376 gtgactgata taaaagagaa aaaaggcgat tttccgcct aaaactctt aaaacttatt 6720
 E--> 1378 aaaactctta aaaccggcct ggcctgtcga taactgtctg gccagcgcac agccgaagag 6780
 E--> 1380 ctgcaaaaag cgccttaccct tcggtcgtc cgctccctac gccccggc ttcgcgtcgg 6840
 E--> 1382 cctatcgccg cgcgtggccg ctcaaaaatg gctggcctac ggccaggcaa tctaccagg 6900
 E--> 1384 cgcggacaag cgcggccgtc gccactcgac cgccggcgct gaggtctgcc tcgtgaagaa 6960
 E--> 1386 ggtgttgcgtc actcatacca ggcctgaatc gcccattat ccagccagaa agtgagggag 7020
 E--> 1388 ccacgggtga tgagagctt gtttaggtg gaccaggttt tgattttgaa cttttgctt 7080
 E--> 1390 gccacggaaac ggtctgcgtt gtcggagaaga tgctgtatct gatcctcaa ctcagcaaaa 7140
 E--> 1392 gttcgattta ttcaacaaag cgcgttccc gtcaagtca cgtaatgtc tggcagtgtt 7200
 E--> 1394 acaaccaatt aaccaattct gattagaaaa actcatcgag catcaatga aactgcaatt 7260
 E--> 1396 tattcatatc aggattatca ataccatatt tttgaaaaag cggtttctgt aatgaaggag 7320
 E--> 1398 aaaactcacc gaggcagttc cataggatgg caagatcctg gtatcggtc gcgattccg 7380
 E--> 1400 ctctgtccaaac atcaatacaa cctattaatt tccctcgatc aaaataagg ttatcaagt 7440
 E--> 1402 agaaatcacc atgagtgacg actgaatccg gtgagaatgg caaaactctt gatataatga 7500
 E--> 1404 atcgccaaac ggcggggggag aggccgtttt cgtattggc gcttccgc ttcctcgctc 7560
 E--> 1406 actgactcgc tgcgtctcggt cggtcggtc cggcgagccg tatacgctca ctcaaaaggcg 7620
 E--> 1408 gtaatacggt tatccacaga atcaggggat aacgcaggaa agaacatgtg agcaaaaaggc 7680
 E--> 1410 cagcaaaagg ccaggaaccg taaaaaggcc gcgttgcgtt cggtttccaa taggctccgc 7740
 E--> 1412 cccctgacg agcatcacaa aaatcgacgc tcaagtca ggtggcggaaa cccgacagga 7800
 E--> 1414 ctataaagat accaggcggtt tcccccttggaa agtccctcg tgcgtctcc tttccgacc 7860
 E--> 1416 ctgcgcgtt ccggataacct gtccgcctt ctcccttgcgg gaagcggtgc gcttctcat 7920
 E--> 1418 agtcacgct ttaggtatct cagttcggtg taggtcggtc gctccaagct gggctgtgt 7980
 E--> 1420 cacgaacccc ccgttcagcc cgaccgcgtc gccttatccg gtaactatcg tcttgagtcc 8040
 E--> 1422 aacccggtaa gacacgactt atcgccactg gcagcagcca ctggtaacag gattagcaga 8100
 E--> 1424 gcgagggtatg taggcgggtc tacagaggatc ttgaaagtggt ggcttaacta cgctacact 8160
 E--> 1426 agaagaacag tatttggat ctgcgtctcg ctgaagccag ttacctcgg aaaaagagtt 8220
 E--> 1428 ggtagctt gatccggcaa acaaaccacc gctggtagcg gttttttt ttttgcaag 8280
 E--> 1430 cagcagatc cgcgcagaaaa aaaggatct caagcagatc ctttgatctt ttctacgggg 8340
 E--> 1432 Page 21
 W--> 1434 70094 Syngenta *delete*
 E--> 1436 tctgacgcgtc agtggaaacga aaactcacgt taaggattt tggcatgag attatcaaaa 8400
 E--> 1438 aggtatcttca cctagatcct tttgatccgg aattaattcc tgggttggc atgcacatac 8460
 E--> 1440 aaatggacga acggataaac ctttgcacgc cttttaaat atccgattat tctaataaaac 8520
 E--> 1442 gctctttctt cttaggttta cccgccaata tttctgtca aacactgata gttaaactg 8580
 E--> 1444 aaggcggggaa acgacaatct gatcatgagc ggagaattaa gggagtcacg ttatgacccc 8640
 E--> 1446 cgcgcgtac gcccgcgtac ccgttttacg tttggactg acagaaccgc aacgcgtcag 8700
 E--> 1448 gaattggcccg cggccat taaaatcaat tggcgccgc gatttcgagc tcggtagc 8757
 E--> 1450 Page 22

P/S
delete

8755

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 12/22/2004
PATENT APPLICATION: US/10/517,039 TIME: 09:15:17

Input Set : A:\Sequence Listing-70094USPCT.txt
Output Set: N:\CRF4\12222004\J517039.raw

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/517,039

DATE: 12/22/2004
TIME: 09:15:17

Input Set : A:\Sequence Listing-70094USPCT.txt
Output Set: N:\CRF4\12222004\J517039.raw

L:1 M:259 W: Allowed number of lines exceeded, (1) GENERAL INFORMATION:
L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:65 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:1264 SEQ:1
L:65 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:71 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:73 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
M:254 Repeated in SeqNo=1
L:137 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:145 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:147 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:209 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:215 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:217 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:281 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:284 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:286 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:306 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:8
L:312 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:326 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:348 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:352 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:354 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:378 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9555 Found:9575 SEQ:1
L:418 M:254 E: No. of Bases conflict, LENGTH:Input:6 Counted:844 SEQ:2
L:418 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:422 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:424 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
M:254 Repeated in SeqNo=2
L:486 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:489 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:491 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:555 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:557 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:559 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:585 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:589 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:593 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:621 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:625 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:627 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:691 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:694 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:696 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:2
L:734 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9546 Found:9563 SEQ:2
L:758 M:254 E: No. of Bases conflict, LENGTH:Input:11 Counted:364 SEQ:3
L:758 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:760 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/517,039

DATE: 12/22/2004
TIME: 09:15:17

Input Set : A:\Sequence Listing-70094USPCT.txt
Output Set: N:\CRF4\12222004\J517039.raw

L:762 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3
M:254 Repeated in SeqNo=3
L:792 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:826 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:828 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:830 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3
L:892 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:894 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:896 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3
L:914 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:918 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:9
L:920 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:8
L:926 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:928 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:932 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:5
L:940 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:946 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:948 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:960 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:962 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:964 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3
L:1026 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:1029 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:1031 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3
L:1095 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:1098 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:1100 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3
L:1124 M:252 E: No. of Seq. differs, <211> LENGTH:Input:10604 Found:10626 SEQ:3
L:1162 M:254 E: No. of Bases conflict, LENGTH:Input:17 Counted:784 SEQ:4
L:1162 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:1164 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:1166 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:4
M:254 Repeated in SeqNo=4
L:1230 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:1233 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:1235 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:4
L:1235 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:1297 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:1300 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:1302 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:4
L:1366 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:1368 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:1370 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:4
L:1432 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:1434 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:2
L:1436 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:4
L:1450 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:1450 M:252 E: No. of Seq. differs, <211> LENGTH:Input:8757 Found:8779 SEQ:4